#### crt sequences.WorkFile

```
Organization Applicant
         Street: 351 Tache Avenue
         City: Winnipeg
         State : Manitoba
         Country : Canada
         PostalCode: R2H 2A6
         PhoneNumber: 204-235-3206
         FaxNumber:
         EmailAddress:
<110> OrganizationName : St. Boniface General Hospital Research Centre
Individual Applicant
         Street: 351 Tache Avenue
         City: Winnipeg
         State : Manitoba
         Country : Canada
         PostalCode: R2H 2A6 PhoneNumber:
         FaxNumber:
         EmailAddress:
<110> LastName : Mesaeli
<110> FirstName : Nasrin
<110> MiddleInitial :
<110> Suffix :
Application Project
<120> Title : Transgenic Mouse over-expressing calreticulin (CRT) in vascular
           smooth muscle cells
<130> AppFileReference : 81190-2602
<140> CurrentAppNumber :
<141> CurrentFilingDate : ____-__
Earlier Applications
<150> PriorAppNumber : US 60/455399
<151> PriorFilingDate : 2003-03-18
Sequence
<213> OrganismName : Artificial
<400> PresequenceString :
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                                                                                                                 60
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                                                                                                               120
                                                                                                               180
                                                                                                               240
                                                                                                                300
cagatagggg cagaggagag ctggttctgt ctccactgtg tttggtcttg ggtactgaac
                                                                                                                360
                                                                                                               420
tcaqaccatc aggregata gcagttgtct ttaaccctaa ccctgagcct gtctcacctg
tcccttccca agaccactga agctaggtgc aagataagtg gggacccttt ctgaggtggt
                                                                                                                480
aggatette acgataagga etatttgaa gggagggagg gtgacaetgt ectagteete ttaceetagt gteeteeage ettgeeagge ettaaaeate egeecattgt eacegeteta gaaggggeea gggttgaett getgetaaae aaggeaetee etagagaage accegetaga aggataeeat acetgtggge aggatgaeee atgttetgee acgeaettgg tageettgga aaggeeactt tgaaceteaa tttteteaae tgttaaatgg ggtggtaaet getateteat aataaagggg aacgtgaaag gaaggegttt geatagtgee tggttgtgea geeaggetge aggeagaget aggetgeaagget aggetgeaggetge aggatgaeet aggteeagaget aggetgeaggetge
                                                                                                                540
                                                                                                               600
                                                                                                               660
                                                                                                                720
                                                                                                               780
                                                                                                               840
agtcaagact agttcccacc aactcgattt taaagccttg caagaaggtg gcttgttgt cccttgcagg ttcctttgtc gggccaaact ctagaatgcc tccccctttc tttctcattg
                                                                                                               900
                                                                                                               960
aagagcagac ccaagtccgg gtaacaagga agggtttcag ggtcctgccc ataaaaggtt
tttcccggcc gccctcagca ccgccccgcc ccgacccccg cagcatctcc aaagcatgca
gagaatgtct ccggctgccc ccgacagact gctccaaactt ggtgtctttc cccaaatatg
                                                                                                              1020
                                                                                                              1080
                                                                                                              1140
                                                              Page 1
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#### crt sequences.WorkFile gagcctgtgt ggagtgagtg gggcggcccg gggtggtgag ccaagcagac ttccatgggc agggaggggc gccagcggac ggcagagggg tgacatcact gcctaggcgg cctttaaacc cctcaccag ccggcgccc accgagctcg gatccactag tccagtgtgg tggaattcat 1200 1260 1320 gctgctccct gtgccgctgc tgctcggcct gctcggcctg gccgccgccg agcccgtcgt 1380 ctacttcaag gagcagttic tggacggaga tgggtggacc gagcgctgga tcgaatccaa 1440 acacaagtee gattttggea aattegteet eagtteggge aagttetaeg gegateagga 1500 gaaagataaa gggctgcaga ccagccagga cgcccgcttc tacgccctgt cggcccgatt 1560 cgadagatada gggctgcaga ccagccagga cgcccgcttc tacgccctgt cggcccgattc cgagccgttc agcaacaagg gccagccact ggtggtgcag ccagccagga cgcccgcttc tacgccctgt cggcccgatt cgagccgttc agcaacaagg gccagccact ggtggtgcag ttcaccgtga aacacgagca gaacattgac tgcgggggcg gctacgtgaa gctgtttccg gccggcctgg accagaagga catgcacggg gactctgagt acaacatcat gtttggtcct gacatctgtg gccccggcac caagaaggtt cacgtcatct tcaactacaa gggcaagaac gtgctgatca acaaggacat ccgttgcaag gacgacgagt tcacacacct gtacacgctg atcgtgggc cggacaacac gtatgaggtg aagattgaca acagccaggt ggagtcgggc tccctgggagg actgctgaag actggggag actggagaca acaaggaacc agatgcctaga 1620 1680 1740 1800 1860 1920 1980 2040 aagcctgaag actgggacga gcgggccaag atcgacgacc ccacggactc caagcccgag 2100 2160 gactgggacă agcccgagca catccccgac ccggacgcga agaagcccga agactgggac gaagaaatgg acggagatg ggagccgccg gtgattcaga accccgagta caagggtgag tggaagccgc ggcagatcga caaccccgat tacaaaggca cctggatcca ccccgaaatc gacaaccccg agtactcgcc cgacgctaac atctatgcct acgacagctt tgccgtgctg ggcttggacc tctggcaggt caagtcgggc accatcttcg acaacttcct catcaccaac 2220 2280 2340 2400 gatgaggcgt acgcagagga gtttggcaac gagacgtggg gcgtcaccaa gacggccgag 2460 aagčagatga aagacaagca ggacgaggag cagcggatga aggaggagga ggaggagaag 2520

aağcggaagg aggaggagga ggccgaggag gacgaggagg acaaggacga caaggaggac

gaggatgagg acgaggagga caaggacgag gaggaggagg aggcggccgc cggccaggcc

2580

2640

2655

aaggacgagc tgtag
<212> Type : DNA
<211> Length : 2655

SequenceName : SM22a-CRT SequenceDescription :

# Feature

Sequence: SM22a-CRT:
<221> FeatureKey : CDS
<222> LocationFrom : 1319
<222> LocationTo : 2653

Other Information : CRT coding sequence

CDSJoin: No

### Sequence

<213> OrganismName : Artificial						
<400> PreSequenceString :						
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gccaggctgg ctgtagtgga ttgagcgtd	t gaggetgeac	ctctctggcc	tgcagccagt	120		
tcctgggtga gactgaccct gcctgaggg	t tctctccttc	cctctctcta	ctcctttcct	180		
ccctctcct ctccctctct ctgtttcct	g aggtttccag	gattggggat	gggactcaga	240		
gacaccacta aagccttacc ttttaagaa	ig tigcattcag	tgagtgtgtg	agacatagca	300		
cagatagggg cagaggagag ctggttctg	t ctccactgtg	tttggtcttg	ggtactgaac	360		
tcagaccatc aggtgtgata gcagttgto	t ttaaccctaa	ccctgagcct	gtctcacctg	420		
tcccttccca agaccactga agctaggtg	c aagataagtg	gggacccttt	ctgaggtggt	480		
aggatette acgataagga etattttga	aa gggagggagg	gtgacactgt	cctagtcctc	540		
ttaccctagt gtcctccagc cttgccagg	cttaaacatc	cgcccattgt	caccgctcta	600		
qaaqqqqcca qqqttqactt gctgctaaa	ac aaggcactcc	ctagagaagc	acccgctaga	660		
agcataccat acctgtgggc aggatgacc	c atgttctgcc	acgcacttgg	tagccttgga	720		
aaggccactt tgaacctcaa ttttctcaa	ac tgttaaatgg	ggtggtaact	gctatctcat	780		
aataaagggg aacgtgaaag gaaggcgtt	tt gcatagtgcc	tggttgtgca	gccaggctgc	840		
agtcaagact agttcccacc aactcgatt	tt taaagccttg	caagaaggtg	gcttgtttgt	900		
cccttgcagg ttcctttgtc gggccaaac	ct ctagaatgcc	tccccctttc	tttctcattg	960		
aagagcagac ccaagtccgg gtaacaagg	a agggtttcag	ggtcctgccc	ataaaaggtt	1020		
tttcccggcc gccctcagca ccgccccgc	cc ccgacccccg	cagcatctcc	aaagcatgca	1080		
gagaatgict ccggctgccc ccgacagac	ct gctccaactt	ggtgtctttc	cccaaatatg	1140		
	Dago					

Page 2

# crt sequences.WorkFile

gagcctgtgt	ggagtgagtg	gggcggcccg	gggtggtgag	ccaagcagac	ttccatgggc	1200
agggagggc	gccagcggac	ggcagagggg	tgacatcact	gcctaggcgg	cctttaaacc	1260
cctcacccag	ccggcgcccc	accgagctcg	gatccactag	tccagtgtgg	tggaattcat	1320
gctgctccct	gtgccgctgc	tgctcggcct	gctcggcctg	gccgccgccg	agcccgtcgt	1380
ctacttcaag	gagcagtttc	tggacggaga	tgggtggacc	gagcgctgga	tcgaatccaa	1440
acacaagtcc	gattttggca	aattcgtcct	cagttcgggc	aagttctacg	gcgatcagga	1500
		ccagccagga				1560
		gccagccact				1620
		cgagccgttc				1680
		gaacattgac				1740
		catgcacggg				1800
		caagaaggtt				1860
gtgctgatca	acaaggacat	ccgttgcaag	gacgacgagt	tcacacacct	gtacacgctg	1920
		gtatgaggtg				1980
tccctggagg	atgactggga	cttcctaccc	cccaagaaga	taaaggaccc	agatgcctcg	2040
		gcgggccaag				2100
		catccccgac				2160
		ggagccgccg				2220
tggaagccgc	ggcagatcga	caaccccgat	tacaaaggca	cctggatcca	ccccgaaatc	2280
		cgacgctaac				2340
		caagtcgggc				2400
		gtttggcaac				2460
		ggacgaggag				2520
		ggccgaggag				2580
		caaggacgag				2640
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<212> Type : DNA <211> Length : 2691

SequenceName : sm22a-crt-ha

SequenceDescription:

#### Feature

Sequence: sm22a-crt-ha: <221> FeatureKey : CDS <222> LocationFrom : 1319
<222> LocationTo : 2689 Other Information : CDSJoin : No

## Feature

Sequence: sm22a-crt-ha:

<221> FeatureKey : misc\_feature

<222> LocationFrom : 2635
<222> LocationTo : 2670

Other Information : HA insertion CDSJoin : No

# Sequence

<213> OrganismName : Artificial <400> PreSequenceString: MLLPVPLLLG LLGLAAAEPV VYFKEQFLDG DGWTERWIES KHKSDFGKFV LSSGKFYGDQ 60 EKDKGLQTSQ DARFYALSAR FEPFSNKGQP LVVQFTVKHE QNIDCGGGYV KLFPAGLDQK 120 DMHGDSEYNI MFGPDICGPG TKKVHVIFNY KGKNVLINKD IRCKDDEFTH LYTLIVRPDN 180 TYEVKIDNSQ VESGSLEDDW DFLPPKKIKD PDASKPEDWD ERAKIDDPTD SKPEDWDKPE 240 300 HIPDPDAKKP EDWDEEMDGE WEPPVIQNPE YKGEWKPRQI DNPDYKGTWI HPEIDNPEYS PDANIYAYDS FAVLGLDLWQ VKSGTIFDNF LITNDEAYAE EFGNETWGVT KTAEKQMKDK 360 QDEEQRLKEE EEEKKRKEEE EAEEDEEDKD DKEDEDEEE DKDEEEEEAA AGQAKDEL 418

<212> Type : PRT

# crt sequences.WorkFile

<211> Length : 418

SequenceName : crt SequenceDescription :

### Sequence

<pre>&lt;213&gt; OrganismName : Artificial &lt;400&gt; PreSequenceString :</pre>	
A PUBLIC LICEA A PROVENCIA DE DESTERNITES EN CONTROLLES	CCCVEVCDO 60
MLLPVPLLLG LLGLAAAEPV VYFKEQFLDG DGWTERWIES KHKSDFGKFV LS	
EKDKGLQTSQ DARFYALSAR FEPFSNKGQP LVVQFTVKHE QNIDCGGGYV KI	LFPAGLDQK 120
DMHGDSEYNI MFGPDICGPG TKKVHVIFNY KGKNVLINKD IRCKDDEFTH L'	YTLIVRPDN 180
TYEVKIDNSO VESGSLEDDW DFLPPKKIKD PDASKPEDWD ERAKIDDPTD SI	
HIPDPDAKKP EDWDEEMDGE WEPPVIQNPE YKGEWKPRQI DNPDYKGTWI HI	PEIDNPEYS 300
PDANIYAYDS FAVLGLDLWQ VKSGTIFDNF LITNDEAYAE EFGNETWGVT K	
QDEEQRLKEE EEEKKRKEEE EAEEDEEDKD DKEDEDEDEE DKDEEEEEAA AG	
DYARQAKDEL	430
or the state of th	

<212> Type : PRT <211> Length : 430

SequenceName : crtha SequenceDescription :

#### Feature

Sequence: crtha: